

FIG.1

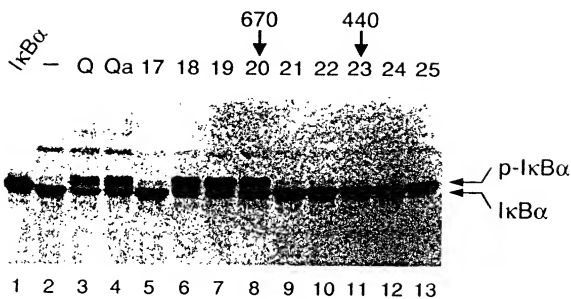


FIG.2A

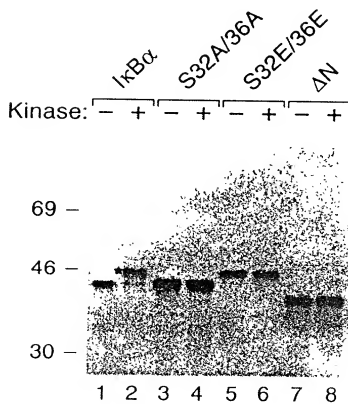


FIG.2C

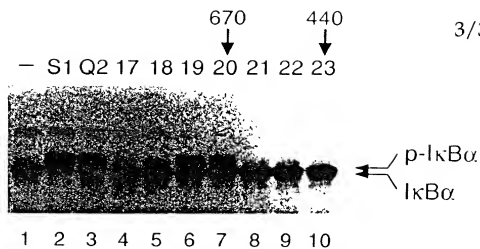


FIG.2B

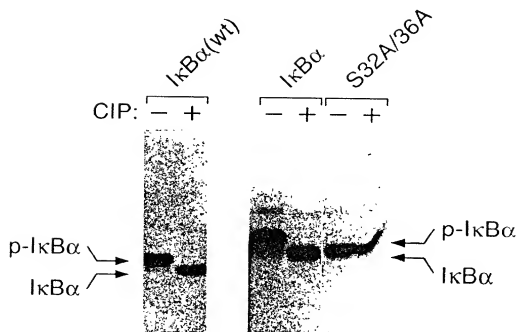


FIG.2D

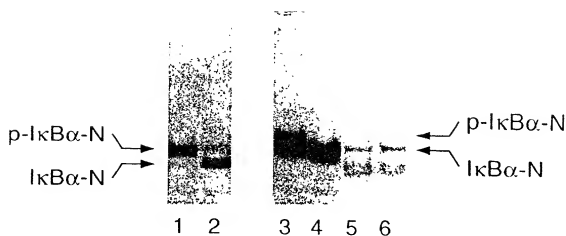


FIG.2E

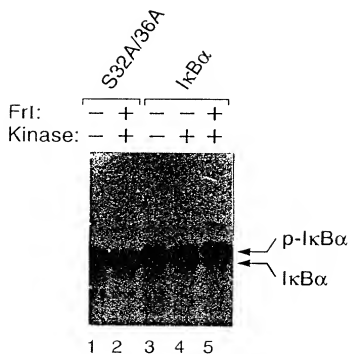


FIG.3A

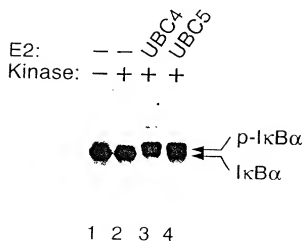


FIG.3B

FIG.3C

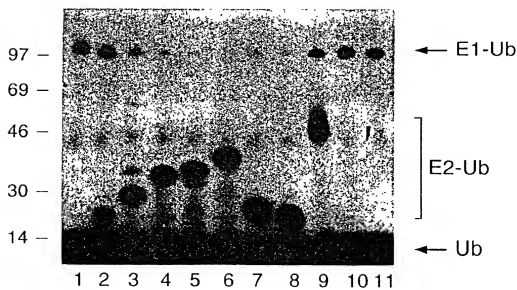
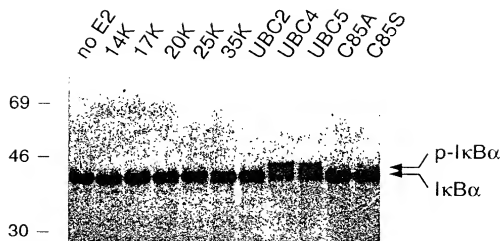


FIG.3D

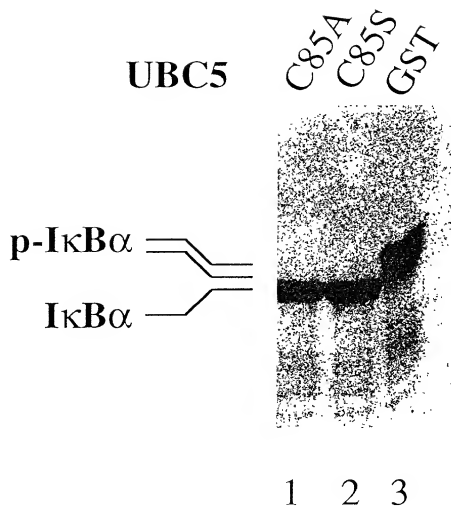


FIG.3E

Ub(μ M): 0 0.1 0.3 1 3 10

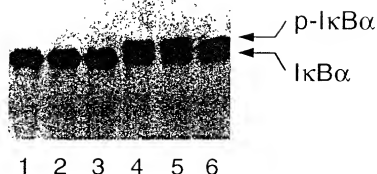


FIG.4A

Order of addition:

Ub(μ M): 0 1.2 2.4 [Ub/MeUb] 2.4 60 0 [MeUb/Ub] 2.4 60

MeUb(μ M): 0 0 0 40 40 40 40 40

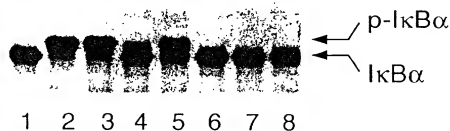


FIG.4B

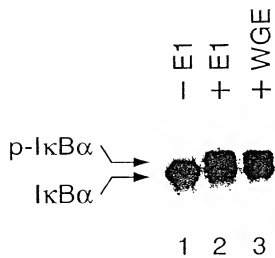


FIG.5A

Kinase: - + + +
 Rel A: + + - +
 OA: + + + -

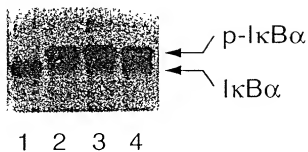


FIG.5B

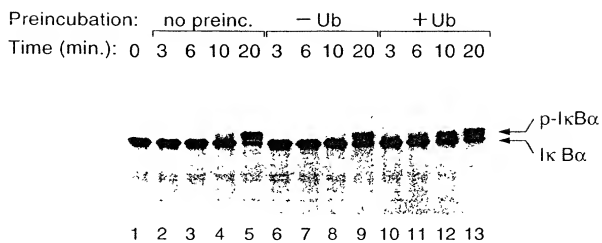


FIG.6A

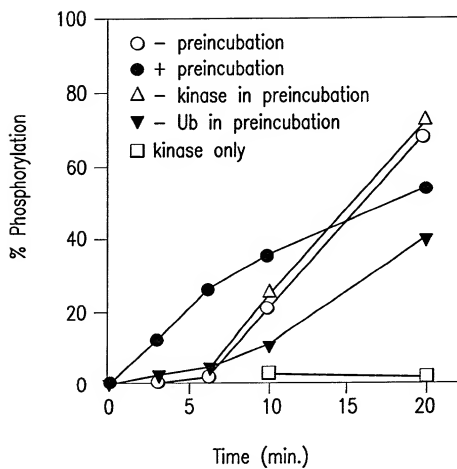


FIG. 6B

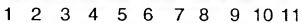


FIG. 7A

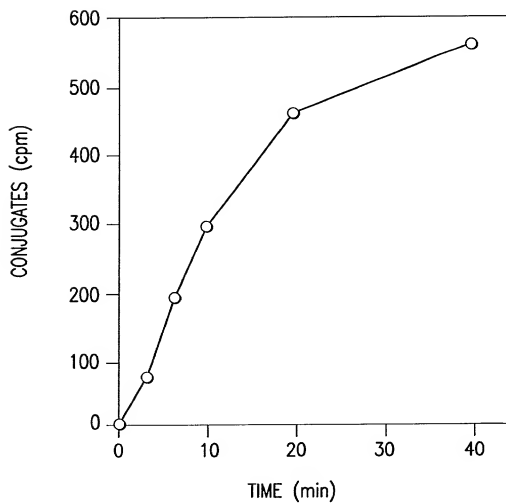


FIG. 7B

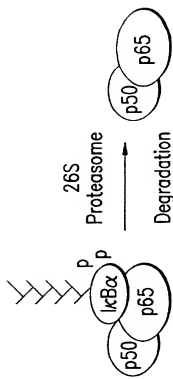
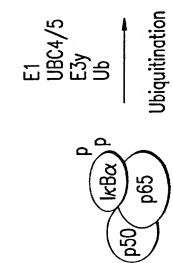
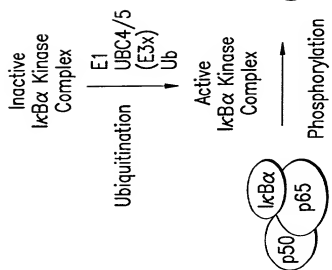


FIG. 8

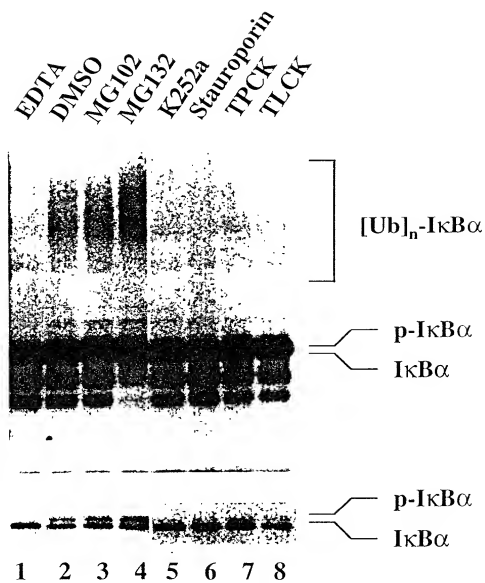


FIG.9

FIG. 10A

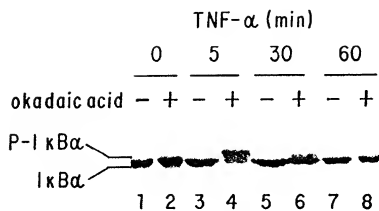
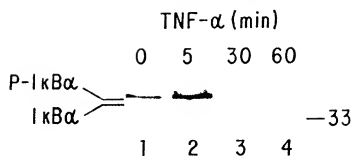


FIG. 10B

FIG. 10C

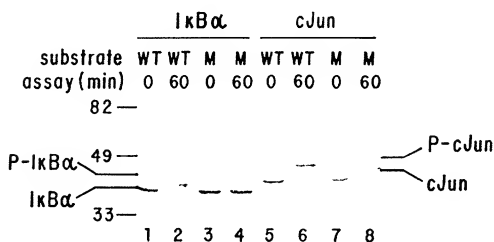
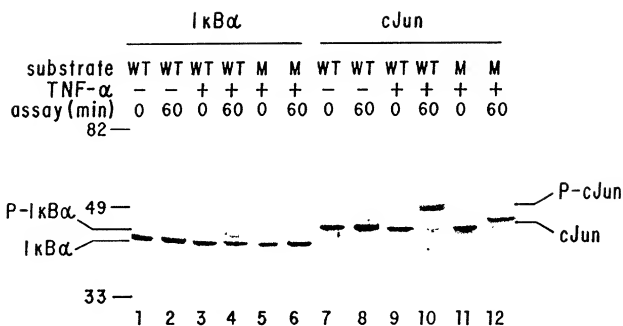


FIG. 10D

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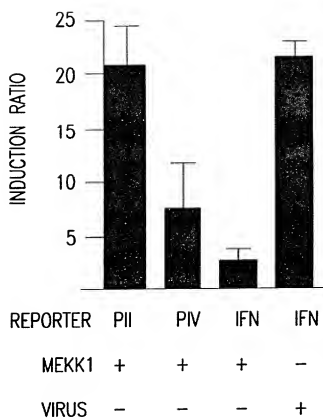


FIG. 11A

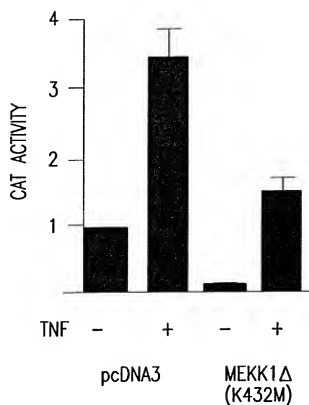


FIG. 11B

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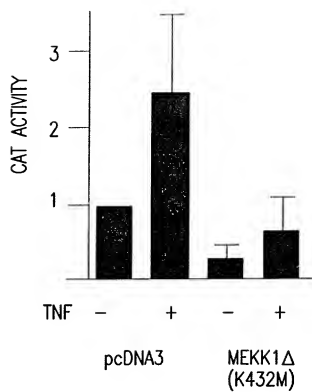


FIG. 11C

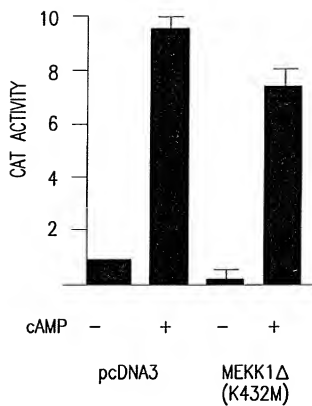


FIG. 11D

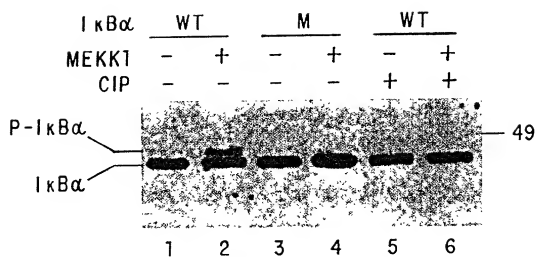


FIG.12

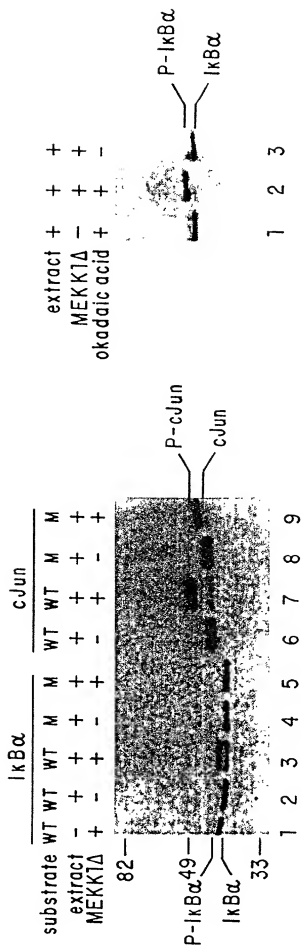


FIG.13A

FIG.13B

FIG. 14A



FIG. 14C



FIG. 14B



FIG. 14D



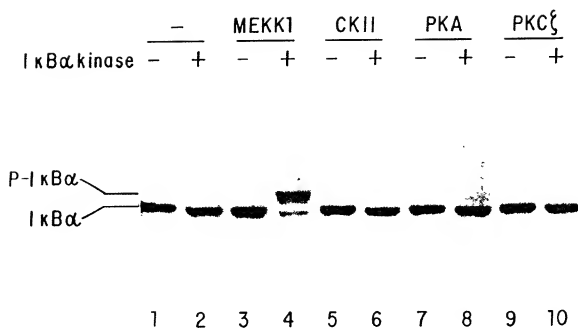


FIG.15A

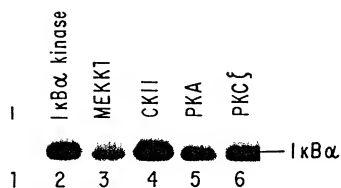


FIG.15B

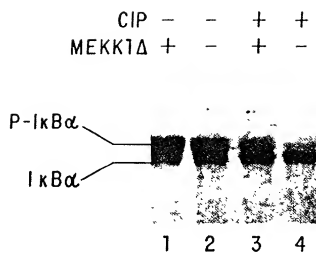


FIG.16A

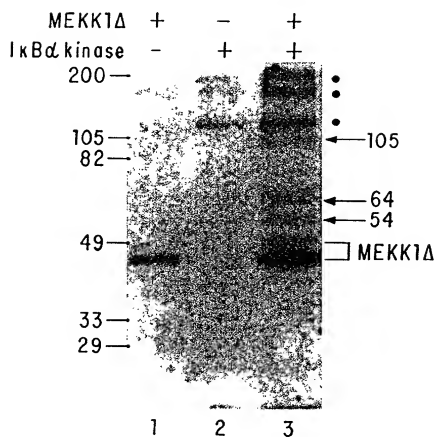


FIG.16B

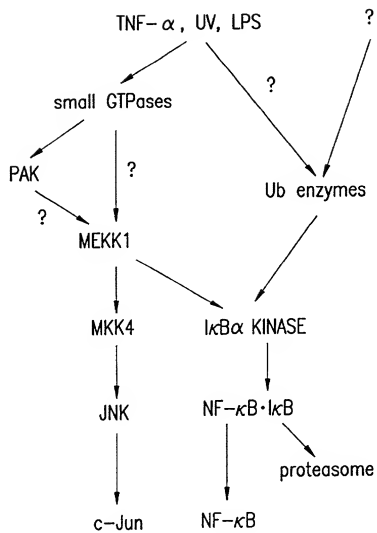


FIG. 17

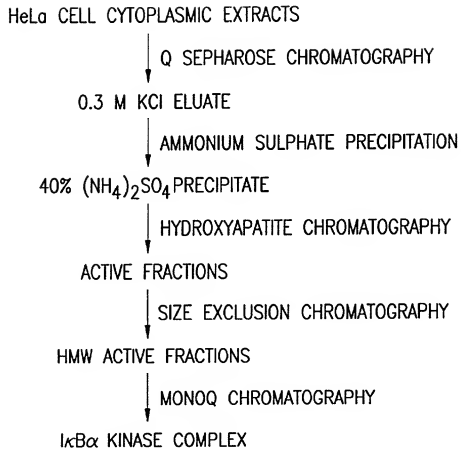


FIG. 18

**Native Gel
(2-15%)**



Hsp90—

FIG.19A

**SDS Gel
(12%)**

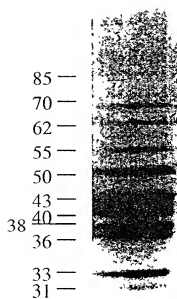


FIG.19B

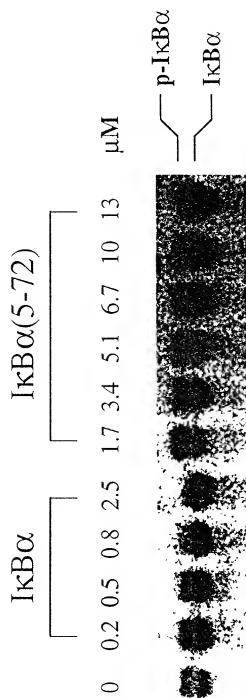


FIG. 20

- A p50: pep1: [L/I] [Y] [V] [E] [L/I] [E] [R]
 B pep2: [L/I] [Q/K] [E] [V] [L/I] [E] [T] [L/I] [L/I] [S]
 [L/I] [E] [K*]
 C pep3: [L/I] [F] [T] [T] [M] [E] [L/I] [M] [R]
 D p40: pep4: [T] [Y] [H] [A] [L*] [S] [N] [L*] [P] [K*]

FIG.21

1 gggtgacgag tgggtggccga agcaggggga cagcaaggga cgctcaggcg gggaccatgg
 61 cggacggcgg ctcgagcgcg gctgacgggc gcatcgtaaa gatggagggtg gactacagcg
 121 ccacgggtgga tcagcgcccta cccgagtggtg cgaagtatgc caaggaaaggga agacttcaag
 181 aagtcattga aacccttctc tctctggaaa agcagactcg tactgcttcc gatatgggat
 241 cgacatcccg tatcttagtt gcagtagtga agntgtgcta tgaggctaaa gaatgggatt
 301 tacttaatta aaaatattat tgctttttgt ccaaaaggcg gagtcaagtt aaaaacaagc
 361 tagttgacaa aaaatggatt naacagttgc tgnacttat tgttt

1 ataccaagag gtaccaggaa gcattgcatt tgggttctca gctgctgcgg gaggatgaaa
 61 agatggacga caaagctctt ttggtggaag tacagctttt agaaagcaaa acataccatg
 121 ccctgagcaa cctgcgaaaa gcccgagctg ccttaacttc ttctcgaacc acagcaaatg
 181 ccatctactg cccctaaaat tgcaggccac cttggacatg cagtcgggta ttatccatgc
 241 agcagaagag aaggcttgaa actcgtactc atacttctat gaggcattta gggatgact
 301 catcgacagc ccaaggcatc aca

FIG.22

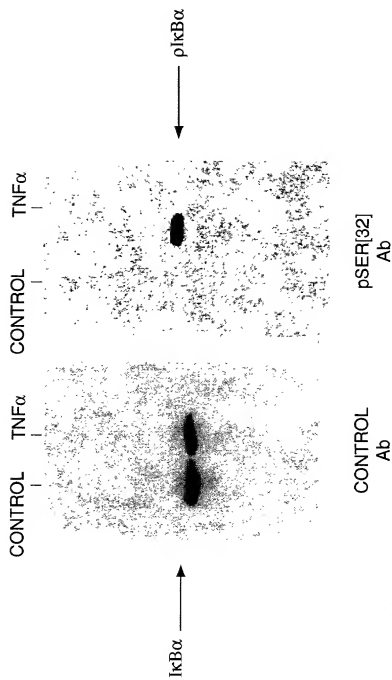


FIG. 23

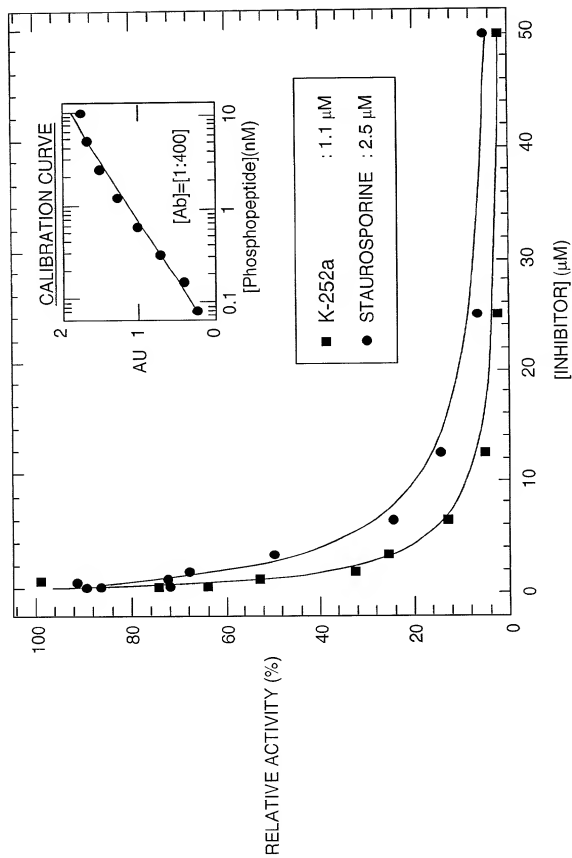


FIG. 24